

CORRECTION

Correction: A Computational Model for Predicting RNase H Domain of Retrovirus

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Reference

1. Wu S, Zhang X, Han J (2016) A Computational Model for Predicting RNase H Domain of Retrovirus. PLoS ONE 11(8): e0161913. doi:[10.1371/journal.pone.0161913](https://doi.org/10.1371/journal.pone.0161913) PMID: [27574780](https://pubmed.ncbi.nlm.nih.gov/27574780/)



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